



FIG S1 Heatmaps of (A) *H. ducreyi* and (B) human genes showing Pearson correlation coefficients (r) between volunteers at infected sites (462, 465, 466, and 467) and (A) inocula or (B) wounded sites (462c, 465c, 466c, and 467c), respectively. Pearson correlation coefficients were calculated after removal of low-expressing genes and transformation of counts to CPM (counts per million).